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1 homo sapien

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2 homo sapien

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9 rattus norv

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1 homo sapien
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RT R	RESU MT 22 LD DT DT DT DT DT DT RESU OCC OCC OCC OCC OCC OCC OCC OCC OCC OCC
"Metallothionein system."; Proc. Natl. Acad. [2] SEQUENCE FROM N.A STRAIN-OREGON-R; MEDLINE-91012582; Silar P., Theodor "Metallothionein regulation."; J. Mol. Biol. 215 SEQUENCE FROM N.A STRAIN-BERKELEY; MEDLINE-20196006; Adams M.D., Celni Amanatides P.G., George R.A., Lewi Sutton G.G., Wort Brandon R.C., Rog Wan K.H., Doyle C Abril J.F., Agbay Ballew R.M., Basu Besson K.Y., Beno Borkova D., Botch Burtis K.C., Busa Cherry J.M., Cawl de Pablos B., Dell Dodson K. J., Evan Fosler C., Gabrie Glodek A., Gong F Harris N.L., Harv Hostin D., Housto Jalali M., Kalush Kimmel B.E., Kodi Liako P., Lei Y., Liu X., Mattei B.	
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A Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,

A Palazzolo M., Pittman G.S., Pan S., Pollard J., Puri V., Reese M.G.,

A Reinert K., Remington K., Sanders R.D.C., Scheeler F., Shen H.,

A Shue B.C., Siden-Klamos I., Simpson M., Skupski M.P., Smith T.,

A Spier E., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,

A Syirskas R., Tector C., Turner R., Venter E., Wang A.H., Wang X.,

Wang Z.-Y., Wassarman D.A., Weinstock G.M., Weissenbach J.,

A Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,

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A Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,

A Williams S.M., Woodage T., Worley K.C., Shan M., Zhang G., Zhao Q., Zheng L.,

A Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L.,

A Zheng X.H., Zhong F.N., Zhong W., Zhou X., Zhu S., Zhu X., Smith H.O.,

RA Gibbs R.A., Myers E.W., Rubin G.M., Venter J.C.,

"The genome sequence of Drosophila melanogaster.",

Science 287:2185-2195(2000).
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Matches
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Ephydroidea; Drosophilidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FLYBASE;
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EMBL; X52098; CAA36318.1; -.
EMBL; AE003730; AAF55768.1;
PIR; S14706; SMFF2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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INDUCTION: STRONGLY INDUCED BY CADMIUM, COPPER AND MERCURY.

INDUCTION: STRONGLY INDUCED BY CADMIUM, COPPER AND MERCURY.

MISCELLANEOUS: ALL CYSTEINE RESIDUES ARE ARRANGED IN C-X-C

THESE ARE THOUGHT TO BE THE METAL-BINDING SITES IN OTHER
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                                                                                                                      Metazoa; Arthropoda;
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PubMed=1684716;
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Matches

Conservative

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Score 68; DB Pred. No. 25; Mismatches

Length 55;

Indels

0;

Gaps

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88.3%; 17.9%;

Best Local Similarity

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RA Harris N.L., Harvey D., Heiman T.J., Hernandez J.R., Houck J.,
RA Hostin D., Houston K.A., Howland T.J., Wein M.-H., Ibeywam C.,
RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Jalali M., Kalush F., Karpen G.H., Ke Z., Kennison J.A., Ketchum K.A.,
RA Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z.,
RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,
RA Liu X., Mattei B., McIntosh T.C., McLeod M.P., McPherson D.,
RA Merkulov G., Milshina N.V., Mobarry C., Morris J., Moshrefi A.,
RA Mentos D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Mount S.M., Moy M., Murphy B., Murphy L., Muzny D.M., Nelson D.L.,
RA Nelson D.R., Nelson K.A., Nixon K., Nusskern D.R., Pacleb J.M.,
RA Reinert K., Remington K., Saunders R.D.C., Scheeler F., Shen H.,
RA Shide B.C., Siden-Kiamos I., Simpson M., Skupski M.P., Smith T.,
RA Spier E., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,
RA Spier E., Spradling A.C., Stapleton M., Skupski M.P., Smith T.,
RA Williams S.M., Woodage T., Weinstock G.M., Weissenbach J.,
RA Williams S.M., Woodage T., Worley K.C., Wu D., Yang S., Yao Q.A.,
RA Ye J., Yeh R.-F., Zaveri J.S., Zhan M., Zhang G., Zhao Q., Zheng L.,
RA Zheng X.H., Zhong F.N., Zhong M., Zhou X., Smith H.O.,
RA Zheng R.A., Myers E.W., Rubin G.M., Venter J.C.;
"The genome sequence of Drosophila melanogaster.";

RT "The genome sequence of Drosophila melanogaster.";
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RA Sutton G.G., Worthan J.R., Yandell M.D., Zhang Q., Chen L.X.,
RA Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Handon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeiffer B.D.,
RA Handon R.C., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,
RA Harris M., Lewis M., Baxter E.G., Helt G., Nelson C.R., Miklos G.L.G.,
RA Harris M., Bayrah A., An H.-J., Andrews-Pfannkoch C., Baldwin D.,
RA Hallew R.M., Bayrah A., Bayraktaroglu L., Beasley E.M.,
RA Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Borkova D., Botchan M.R., Bouck J., Brokstein P., Brottier P.,
RA Cherry J.M., Cawley S., Dahlke C., Davenport L.B., Davies P.,
RA de Pablos B., Delcher A., Deng Z., Mays A.D., Dew I., Dietz S.M.,
RA doson K., Doup L.E., Downes M., Dugan Rocha S., Dunkov B.C., Dunn P.,
RA Durbin K.J. Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W.,
RA Glodek A., Gong F., Gorrell J.H., Gu Z., Guan P., Harris M.,
RA Harris N.I., Hafran T., Gu Z., Guan P., Harris M.,
RA Harris N.I., Hafran T., Harrander I., Harris M.,
FLYBASE; FBgn0004174; Mst84Dc.
Spermatogenesis; Repeat; Multigene family.
SEQUENCE 55 AA; 5225 MW; 95A12F3AEC88BD6C CRC64;
                                                                         EMBL; X67703; CAA47939
EMBL; AE003672; AAF540;
HSSP; P01180; 1NPO
                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "A cluster of row, of Drosophila melanogaster.

of Drosophila melanogaster.

- Dev. 35:143-151(1991).
                                                                                                                                                                        modified and this statement is not removed entitles requires a license agreement (See or send an email to license@isb-sib.ch).
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Adams M.D., Celniker S.E., Holt R.A., Ex
Amanatides P.G., Scherer S.E., Li P.W.,
                                                                                                                                                                                                                                                                                                                                                                                                                             -!- TISSUE SPECIFICITY: TESTIS.
-!- DEVELOPMENTAL STAGE: PRIMARY SPERMATOCYTES.
-!- DOMAIN: THIS PROTEIN IS MOSTLY COMPOSED OF REPETITIVE C-G-P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN-BERKELEY
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Schaefer M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                 SIMILARITY: BELONGS TO THE MST(3)CGP FAMILY.
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                                                                                                 AA47939.1; -.
AAF54025.1;
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                                                                                                                                                                                                                                                       as its content
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                                                                                                                                                                                                                          Usage
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.A., Galle
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MRESULTY 3
MR4A_DROME
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RX 
                                                                                   RA Adams N.D., Celniker S.E., Holt R.A., Evans C.A., Gocayne J.D., RA Adams M.D., Celniker S.E., Holt R.A., Hoskins R.A., Galle R.F., RA Amanatides P.G., Scherer S.E., Li P.M., Hoskins R.A., Galle R.F., RA Amanatides P.G., Scherer S.E., Edhards S., Ashburner M., Henderson S.N., Sutton G.G., Wortman J.R., Yandell M.D., Zhang Q., Chen L.X., RA Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeliffer B.D., Brandon R.C., Rogers Y.-H.C., Blazej R.G., Champe M., Pfeliffer B.D., Baltwin M., An H.-J., Andrews-Pfannkoch C., Baldwin D., Ballwin M., An H.-J., Andrews-Pfannkoch C., Baldwin D., Ballwin R.M., Basu A., Baxendale J., Bayraktaroglu L., Beasley E.M., Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Ballwin K.C., Busam D.A., Butler H., Cadieu E., Center A., Chandra I., Beeson K.Y., Benos P.V., Berman B.P., Bhandari D., Bolshakov S., Burtis K.C., Busam D.A., Butler H., Cadieu E., Center A., Dietz S.M., Beeson K.Y., Dup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., Bodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., Bodson K., Doup L.E., Downes M., Dugan-Rocha S., Dunkov B.C., Dunn P., Dowles R., Dourbin K.J., Evangelista C.C., Ferraz C., Ferriera S., Fleischmann W., Rosler C., Gabrielian A.E., Garg N.S., Gelbart W.M., Glasser K., Ra Durbin K.J., Barvey D., Heiman T.J., Hernandez J.R., Houck J., Harris M.L., Harvey D., Heiman T.J., Wei M.-H., Ibegwam C., Ra Hostin D., Houston K.A., Howland T.J., Wei M.-H., Ibegwam C., Ra Hostin D., Houston K.A., Nowland T.J., Wei M.-H., Ibegwam C., Ra Hostin D., Houston K.A., Nixon K., Wilson J.A., Ketchum K.A., Kimmel B.E., Kodira C.D., Kraft C., Kravitz S., Kulp D., Lai Z., Lin X., Li J., Ling Y., Lin X., Li J., Ling Y., Lin X., Li J., Ling Y., Lin X., Liu X., Mattei B., McIntosh T.C., Morris J., Moshrefi A., Rheriston D.R., Nealson D.L., Rosley B.M., Nelson D.L., Rosley B.M., Nelson D.L., Rosley B.M., Nelson D.L., Rosley B.M., Nelson D.R., Shupski M.P., Smith T., Shen H., Wang X., Wang A.H., Wang X., Wang A.H., Wang X., Wang A.H
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Q01642; Q9VIA3;
Q1-JUL-1993 (Re)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation updat)
MALE SPECIFIC SPERM PROTEIN MST84DA.
                                                                         -!- DEVELOPMENTAL STAGE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=20196006; PubMed=10731132;
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Pterygota;
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ch. Dev. 35:143-151(1991).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Neoptera;
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                                      PROTEIN
                                      IS MOSTLY
                                                                         PRIMARY SPERMATOCYTES
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                                      COMPOSED OF
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                                      REPETITIVE C-G-P
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01-JUL-1993 (Rel. 26,
01-NOV-1997 (Rel. 35,
MALE SPECIFIC SPERM I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              M84D_DROME
Q01645;
                                                                              modified and this statement is not removed entities requires a license agreement (See or send an email to license@isb-sib.ch).
                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboratic between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it
                                                                                                                                                                                                                                            of Drosophila melanogaster.";
Mech. Dev. 35:143-151(1991).
-i- TISSUE SPECIFICITY: TESTIS
                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Musc
Pphydroidea; Drosophilidae; Drosophila.
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use by non-profit institutions as long as its content
modified and this statement is not removed. Usage by ar
 Spermatogenesis; Repeat; SEQUENCE 68 AA; 6480
                           FLYBASE; FBgn0004175; Mst84Dd
                                                                                                                                                                                                                  -!- DEVELOPMENTAL STAGE: PRIMARY SPERMATOCYTES.
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FLYBASE; FBgn0004172; Mst84Da.
                                                      EMBL; X67703;
                                                                                                                                                                                         -!- SIMILARITY: BELONGS TO THE MST(3)CGP FAMILY.
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                                                                                                                                                                                                                                                                                                                    Kuhn
                                                                                                                                                                                                                                                                                                                                MEDLINE=92102953; PubMed=1684716;
                                                                                                                                                                                                                                                                                                                                                STRAIN=OREGON-R;
                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                              Drosophila melanogaster (Fruit fly)
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                                                                                                                         non-profit institutions as long
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                                                      CAA47940.1;
                                          1NPO
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26, Last sequence update)
35, Last annotation updat
RM PROTEIN MST84DD.
                                                                                                                                                                                                                                                                                                                   Boersch
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; Multigene family.
MW; 2F2BD26128DE3DEF
                                                                                                                                                                                                                                                                                                                  D., Glaetzer K.H.,
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                                                                                                                                                                                                                                                                                                                  Schaefer
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Query Match
Best Local Similarity
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Score 68; Pred. No.

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                                                                                                                                                                           01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation updat)
WDNM1 PROTEIN PRECURSOR.
STRAIN-FVB/N;
                                                                          Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                              Mus musculus (Mouse)
                                                                                                                                                                                                                                                                                            Q62477;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FLYBASE; FBgn0004173; Mst84Db.
Spermatogenesis; Repeat; Multigene family.
SEQUENCE 74 AA; 6824 MW; D0793137A7E7D1D0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboratic between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use by non-profit institutions as long as its content is in no was modified and this statement is not removed. Usage by and for commercial modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "A cluster of four genes selectively expressed of Drosophila melanogaster."; Mech. Dev. 35:143-151(1991).
-i- TISSUE SPECIFICITY: TESTIS.
-i- DEVELOPMENTAL STAGE: PRIMARY SPERMATOCYTES.
-i- DOMAIN: THIS PROTEIN IS MOSTLY COMPOSED OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q01643;
01-JUL-1993
                      SEQUENCE
                                                                                                                                                                                                                                                                                                                       WDNM_MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; X67703; CAA47938.1; -. HSSP; P01180; 1NPO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Drosophila melanogaster (Fruit fly).
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Musc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JUL-1993
01-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Schaefer M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=92102953; PubMed=1684716;
Kuhn R., Kuhn C., Boersch D., Glaetzer K.H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRAIN=OREGON-R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ephydroidea; Drosophilidae; Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  M84B_DROME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SIMILARITY: BELONGS TO THE MST(3)CGP FAMILY.
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OV-1997 (Rel. 35, Last annotation update)
SPECIFIC SPERM PROTEIN MST84DB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         CGSCCSPCGSCCAPWGPCGPCGPCCGGC
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N.A.
TISSUE=BREAST
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                                                                          Chordata;
Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 68;
Pred. No.
                                                                       Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
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RESULT
WDNM_RAT
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Best Local Similarity 17.9%;
Matches 5; Conservative
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01-APR-1990 (Rel. 1
01-NOV-1997 (Rel. 3
30-MAY-2000 (Rel. 3
                                                      "The WDNM1 gene product is a novél member of the 'four-disulphide core' family of proteins.";
Biochem. Biophys. Res. Commun. 176:247-254(1991).
-!- FUNCTION: INVOLVED IN THE METASTATIC POTENTIAL OF ADENOCARCING IN RAT. COULD HAVE PROTEINASE INHIBITING CAPACITY.
-!- TISSUE SPECIFICITY: SPLEEN, KIDNEY, BRAIN, LIVER, LUNG, AND CI
                                                                                                                                                 MEDLINE=91207400; PubMed=2018519; Dear T.N., Kefford R.F.;
                                                                                                                                                                                                                                      Dear T.N., Ramshaw I.A., Kefford R.F.; "Differential expression of a novel gemanmary adenocarcinoma cells.";
                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
TISSUE=BREAST CARCINOMA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
This SWISS-PROT entry is copyright.
                           -!- SIMILARITY: BELONGS TO THE WAP-TYPE 'FOUR-DISULFIDE CORE' FAMILY
                                                                                                                                                                               TISSUE=THYMUS;
                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                        Cancer Res. 48:5203-5209(1988).
                                                                                                                                                                                                                                                                                 MEDLINE=88310901; PubMed=3136918;
                                                                                                                                                                                                                                                                                                                                            Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                          Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Morrison B.W., Leder P.;
"neu and ras initiate murine mammary tumors that share genetic markers generally absent in c-myc and int-2-initiated tumors.";
Oncogene 9:3417-3426(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WDNM_RAT
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PROSITE; PS00317; 4_DISULFIDE_CORE;
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35, Last sequence up
39, Last annotation
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                                                                                                                                                                                                                                                                                                                                         Chordata; Craniata; Vertebrata; Euteleostomi; Rodentia; Sciurognathi; Muridae; Murinae; Rat
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Pred. No.
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produced through a collaboration
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RESULT 6
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                                                                                                                                                                                                      the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                        Submitted (MAY-1995) to the EMBL/GenBank/DDBJ databases :- FUNCTION: POSSIBLE FUNCTION IN SPERM MATURATION .- SUBCELLULAR LOCATION: SECRETED (POTENTIAL).
-- TISSUE SPECIFICITY: EPIDIDYMIS.
                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
MAJOR EPIDIDYMIS-SPECIFIC PROTEIN E4 (EPIDIDYMAL PROTEIN BE-20)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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HSSP; P19957; 2REL.
                                                                                              INTERFAC, ---
PFAM; PF00095; wap; 2.
PFAM; PF000917; 4_DISULFIDE_CORE;
                                                                                                                                             EMBL; U26725; AAA66525.1; HSSP; P19957; 1FLE.
                                                                                                                                                                              entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch)
                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                                                                                                                        TISSUE=EPIDIDYMIS; Wang L.F.;
                                                                                                                                                                                                                                                                                                                                                                                                 Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q28631;
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                                              CARBOHYD
                                                             DOMAIN
                                                                                                                                 INTERPRO;
                                                                                                                                                                                                                                                                            -!- TISSUE SPECIFICITY: EPIDIDYMIS.
                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                          Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                             (FRAGMENT)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           7740 MW;
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Pred. No.
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WDNM1 PROTEIN.
51802C70CDAF0521 CRC64;
                                                        WAP 1.
WAP 2.
 Score
                                   N-LINKED (GLCNAC
58BAF723BD74D7F3
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DB
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                                   CRC64;
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Length
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RESULT 10
GAS1_A
ID GAS2_A
AC P46689
DT 01-NOV
DT 15-DEC
DE GIBBER
GN GASA1.
OS ARABIA
OC EUKARY
                                                                                                                                                                            밁
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RSI1_LYCES
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Best Local S
Matches 5
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Matches 5; Conservative
                                                   GAS1_ARATH
P46689;
01-NOV-1995
01-NOV-1995
15-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RS11_LYCES
P47926;
01-FEB-1996
01-FEB-1996
01-NOV-1997
                                                                                                                                                                                                                                                                                                                     Signal.
                                         01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
GIBBERELLIN-REGULATED PROTEIN 1 PRECURSOR.
                Arabidopsis
                                                                                                                                                                                                                                                                                                           CHAIN
                                                                                                                                                                                                                                                                                                                                                                                       or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                               use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MO1. Gen. Genet. 243:148-157(1994).
-!- SUBCELLULAR LOCATION: SECRETED (POTENTIAL).
-!- TISSUE SPECIFICITY: EXPRESSED VERY EARLY IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Taylor B.H., Scheuring C.F.;
"A molecular marker for lateral root initiation: the RSI-1 gene tomato (Lycopersicon esculentum Mill) is activated in early late root primordia.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lycopersicon esculentum (Tomato).
Lycopersicon esculentum (Tomato),
Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
Magnoliophyta; eudicotyledons; core eudicots; Asteridae; euasterids I;
Solanales; Solanaceae; Solanum.
  Eukaryota;
                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                   EMBL;
                                                                                                                                                                                                                                                                                                                                                             EMBL; L22188;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=CV. VFNT; TISSUE=ROOT; MEDLINE=94232181; PubMed=8177211;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RSI-1 PROTEIN PRECURSOR (TR132). RSI-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                         the European Bioinformatics Institute.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- PTM: SIX DISULFIDE BONDS MAY BE PRESENT.
-!- SIMILARITY: BELONGS TO THE GAST1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -!- INDUCTION:
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                                                                                                                                                                                                                                                                                                                                                  L22189;
                                                                                                                                                                                                                              Similarity 17.9
5; Conservative
  Viridiplantae;
                                                                                                                                                                                                                                                                                               96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (Rel. 33, Created)
(Rel. 33, Last sequence up
(Rel. 35, Last annotation
               thaliana
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                                                                                                                                                                                                                                                                                                                                                  AAA20130.1;
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                                                                                                           STANDARD;
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96
                (Mouse-ear cress).
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 Embryophyta;
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                                                                                                                                                                                                                                          Score 68;
Pred. No.
                                                                                                                                                                                                                                                                                           POTENTIAL.
RSI-1 PROTEIN.
C17C826648148CE3 CRC64;
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Mismatches
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Tracheophyta; Spermatophyta;
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                                                                                                                                                                                                                                                        DB 1;
                                                                                                           AA.
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RESULT 11
GAS2_ARATH
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                              Plant
                                                                                                   STRAIN=CV. COLUMBIA;
MEDLINE=95244835; Pub
                                                                                                                                 SEQUENCE FROM
                                                                                                                                                                Brassicales;
                                                                                                                                                                                           Eukaryota; Viridipiantae;
                                                                                                                                                                                                         Arabidopsis thaliana (Mouse-ear cress).
                                                                                                                                                                                                                         GASA2
                                                                                                                                                                                                                                    GIBBERELLIN-
                                                                                                                                                                                                                                                   01-NOV-1995
01-FEB-1996
                                                                                                                                                                                                                                                                                  01-NOV-1995
                                                                                                                                                                                                                                                                                                P46688;
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EMBL; Z29940;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=CV. COLUMBIA; TISSUE=SILIQUES; MEDILINE=95244835; PubMed=7727751; Herzog M., Dorne A.-M., Grellet F.; "GASA, a gibberellin-regulated gene family related to the tomato GAST1 gene."; Plant Mol. Biol. 27:743-752(1995).
                                                        related to
                                                                                     Herzog M.,
                                                                                                                                                                            Magnoliophyta;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      or send
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Multigene
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STEPS OF GEI
SUBCELLULAR
                                       s SWISS-PROT entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the European Bioinformatics Institute. There are no restroy non-profit institutions as long as its content
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TISSUE SPECIFICITY: EXPRESSED IN FLOWER
LOWER LEVELS SEEN IN THE ROOTS AND ROSET
                            FUNCTION: INVOLVED IN LATE STAGES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FUNCTION: MAY FUNCTION IN SOME DEVELOPMENT.
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requires a license agreement (See
an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CTION: BY GIBBERELLINS.
SIX DISULFIDE BONDS MAY BE PRESENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            family;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98
                                                                                                                                                                                                                                    (Rel. 32,
(Rel. 32,
(Rel. 33,
-REGULATED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    57-98 FROM N.A.
               GERMINATION
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                                                                                                                                                              Brassicaceae;
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CAA82842.1;
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                                                                                                                                                                          eudicotyledons; core eu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              eudicotyledons; core eudicots; Rosidae; eurosids II;
                                                                                                                                                                                                                                                                                                             STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98
10718 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         88.3%;
17.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ., Laudie M., Meyer Y., Cooke R., Delseny to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                  Created)
                                                                                                                                                                                                                                                  Last sequence update)
Last annotation updat
                                                                                                                                                                                                                                      PROTEIN 2 PRECURSOR.
                                                                                                                                                              Arabidopsis
                                                                                                                                                                                                                                                                                                                                                                                                                                             23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 68;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GIBBERELLIN-REGULATED PROTEIN BB37B5F0A7C6C89B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               POTENTIAL
                                                                                                                                                                                                                                                                                                             PRT;
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                                                                      family
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                            OF
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                                                                                                                                                                                        Tracheophyta; Spermatophyta;
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37;
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                            SEED MATURATION, OR IN EARLY
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                                                                       thaliana
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밁 20

Query Match Best Local Similarity

88.3%; 17.9%;

Score Pred.

68; No.

DВ 37;

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Length

99

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RESULT
GAS3_AF
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Best Local S
Matches 5
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3_ARATH
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P46687;
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                   01-NOV-1995 (Rel.
01-NOV-1995 (Rel.
01-FEB-1996 (Rel.
                                                                                                                           This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN the European Bioinformatics Institute. There are no restr
                          SIGNAL
                                                                                         entities
                                                                                                                                                                                                                               Herzog M., Dorne A.-M., Grellet F.; "GASA, a gibberellin-regulated gene "related to the tomato GAST1 gene."; Plant Mol. Biol. 27:743-752(1995).
-i- FUNCTION: INVOLVED IN LATE STAGE STEPS OF GERMINATION.
                                                                                                                                                                                                                                                                                                         STRAIN=CV. COLUMBIA; TISSUE=SEED; MEDLINE=95244835; PubMed=772,7751;
                                                                                                                                                                                                                                                                                                                                                      Magnoliophyta; euarcucyrum
mrassicales; Brassicaceae;
                                                                                                                                                                                                                                                                                                                                                                                                              GASA3
                                                                                                                                                                                                                                                                                                                                                                                                                          GIBBERELLIN-REGULATED
 SEQUENCE
                                      Multigene
                                                  EMBL; U11764; AAB06308.1;
                                                                            or send an
                                                                                                    modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                               Arabidopsis thaliana (Mouse-ear cress).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SIGNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Multigene
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                                                                                                                                                                          TISSUE SPECIFICITY: SILIQUES AND DRY SEE PTM: SIX DISULFIDE BONDS MAY BE PRESENT. SIMILARITY: BELONGS TO THE GAST1 FAMILY.
                                                                                                                                                                                                                   SUBCELLULAR LOCATION: SECRETED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SIMILARITY: BELONGS TO THE GAST1 FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE SPECIFICITY: DRY SEEDS AND MATURATING PTM: SIX DISULFIDE BONDS MAY BE PRESENT.
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5; Conser
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                                                                                       requires
                                                                                                                non-profit
                                     family;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       family;
 99
                                                                          email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                       Viridiplantae; Embryophyta; Trachecyta; eudicotyledons; core eudicots;
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              19
 ΑA;
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                                   Signal.
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99
10531 MW;
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17.9%;
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Last annotation updat
PROTEIN 3 PRECURSOR.
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 ¥
                                                                                                                                                                                                                                                                                                                                                             Arabidopsis
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POTENTIAL.
GIBBERELLIN-REGULATED PROTEIN
84549B74B08A47FA CRC64;
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Pred.
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GIBBERELLIN-REGULATED
63713C986887573B CRC6
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                                                                                                                                                                                                                                              STAGES
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                                                                                                                                                                                                                                                                                                                                                                       Tracheophyta; Spermatophyta;
dicots; Rosidae; eurosids II;
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37;
                                                                                                               There are no restrictions ng as its content is in
                                                                                                                                                                                                                                              SEED
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                                                                                     http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                from Arabidopsis thaliana
                                                                                                    Usage
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Matches

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P46690;

THE SPECIES

'FOUR-DISULFIDE CORE'

FROM

EGG

WHITE,

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Query Match
Best Local S
Matches 5
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P00993;
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01-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 use by non-profit institutions as long as its content modified and this statement is not removed. Usage by an entitles requires a license agreement (See http://www.ir.
                                                                                                                                              21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
CHELONIANIN (BASIC PROTEASE INHIBITOR) (RTPI).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Merzog M., Dorne A.-M., Grellet F.; "GASA, a gibberellin-regulated generaled to the tomato GAST1 gene.", Plant Mol. Biol. 27:743-752(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-1995 (Rel. 32, Last sequence update) 01-NOV-1995 (Rel. 32, Last annotation updat GIBBERELLIN-REGULATED PROTEIN 4 PRECURSOR.
                                                                                                   Caretta caretta (Loggerhead).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -!- SUBCELLULAR LOCATION: SECRETED. '
-!- TISSUE SPECIFICITY: DEVELOPING ROOTS
-!- INDUCTION: BY GIBBERELLINS.
    TISSUE=EGG WHITE;
Kato I., Tominaga
                                                                                    Testudines;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Multigene family; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; U15683; AAA74480.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=CV. C24; T: MEDLINE=95244835;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Brassicales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Viridiplantae; Embryophyta; Magnoliophyta; eudicotyledons; core euc
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                                                                                                                                                                                                                                                                                                                                                       46 TQCPSECDRRCKKTQYHKACITFCNKCC
                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
les 5; Conser
                                                                                                                                                                                                                                                                                                                                                                                           SIMILARITY: BELONGS TO THE GAST1 FAMILY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26
106 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Rel.
(Rel.
(Rel.
                                                                                  Cryptodira; Chelonioidea; Cheloniidae; Caretta
                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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5; PubMed=7727751;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              32, Created)
32, Last sequences
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12065 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 68; DB
Pred. No. 39;
23; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     POTENTIAL.
GIBBERELLIN-REGULATED PROTEIN 4.
; 62A3E4A820F57678 CRC64;
                                                                                                                                                                                                                                                   PRT;
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39;
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    RESULT 15
GST1_LYCES
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Best Local S
Matches 5
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01-AUG-1992
01-FEB-1996
                                 This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Extended 
                                                                                                                                                                                                     -!- SUBCELLULAR LOCATION: SECRETED.
-!- TISSUE SPECIFICITY: ALL SHOOT ORGANS.
-!- INDUCTION: BY GIBBERELLINS. INHIBITED
                                                                                                                                                                                                                                                                  from tomato.";
Plant J. 2:153-159(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOD_RES
ACT_SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PFAM; PF00014; Kunitz_BPTI; 1. PFAM; PF00095; wap; 1. PRINTS; PR00003; 4DLSULPHCORE. PRINTS; PR00759; BASICPTASE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TURTLE", THE SPE
PIR; A01224; TITTOR.
HSSP; P00974; 1BRB.
entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                           MEDLINE=93251105; PubMed=1302047;
Shi L., Gast R.T., Gopalraj M., Olszewski N.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lycopersicon esculentum (Tomato) Eukaryota; Viridiplantae; Embryo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GAST1
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                                                                                                                                                               -1- PTM: SIX DISULFIDE BONDS MAY BE PRESENT.
-1- SIMILARITY: BELONGS TO THE GAST1 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                          STRAIN=CV.
                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Viridiplantae; Embryophyta; Tracheophyta; Spermatophyta;
Magnoliophyta; eudicotyledons; core eudicots; Asteridae; euasterids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROSITE; PS00280; BPTI_KUNITZ_1; PROSITE; PS50279; BPTI_KUNITZ_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OF TWO NONHOMOLOGOUS DOMAINS.
-!- SIMILARITY: THE FIRST DOMAIN BELONGS TO THE BPTI/KUNITZ FAMILY OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fed
                                                                                                                                                                                                                                                                                                                                                                                                                                 Solanales;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INTERPRO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -
                                                                                                                                                                                                                                                                                                          "Characterization of a shoot-specific;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INTERPRO; IPR002221;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 XXCXXXCXXXXXXXXXXXXXXXXXXXXXXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INHIBITORS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROTEIN
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5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                      MONEYMAKER; TISSUE-SHOOT;
                                                                                                                                                                                                                                                                                                                                                                                                                               Solanaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      38:832-832(1979).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            110 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (Rel. 23, Created)
(Rel. 23, Last sequence up
(Rel. 33, Last annotation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AS
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Pred. No. 40;
23; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INHIBIT TRYPSIN.
INHIBIT SUBTILISIN.
PYRROLIDONE CARBOXYLIC ACID.
REACTIVE BOND (TRYPSIN).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         update)
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                     (See http://www.isb-sib.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    update)
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